

Molecular Biology  
A SELEX PROCEDURE FOR DETERMINING OPTIMAL DNA BINDING SITES OF  
THE *Pseudomonas aeruginosa* REGULATORY PROTEIN AlgZ

Jenni E. Crowley, Patricia J. Baynham, Ph.D.<sup>\*</sup>, and Daniel J. Wozniak, Ph.D.<sup>‡</sup>

<sup>\*</sup>Thomas More College, 333 Thomas More Parkway, Crestview Hills, KY 41017,  
[trish.baynham@thomasmore.edu](mailto:trish.baynham@thomasmore.edu)

<sup>‡</sup>Department of Microbiology and Immunology, Bowman Gray School of Medicine,  
Wake Forest University, Winston-Salem, NC, 27157-1064

*Pseudomonas aeruginosa* is a gram-negative bacterium found in soil and water. It is an opportunistic pathogen in patients with the genetic disorder cystic fibrosis (CF). *P. aeruginosa* initially colonizes the lungs of CF patients with nonmucoid strains, but these strains later mutate, secrete the exopolysaccharide alginate, and express a mucoid (slimy) phenotype. Isolation of mucoid colonies from the CF lung is associated with a poor prognosis for the patient because alginate provides resistance to the human immune system.

Transcriptional activation of the *P. aeruginosa* alginate biosynthetic operon results in the synthesis of alginate. *algD* is the first gene of this operon and encodes GDP-mannose dehydrogenase, an enzyme that commits precursor molecules to the production of alginate. The regulatory protein AlgZ is crucial for alginate production as both an activator of *algD* transcription and also a repressor of its own synthesis. The sequences of the two known binding sites of AlgZ to the genome of *P. aeruginosa* differ greatly, although a centralized motif has been hypothesized. A dSELEX (systematic evolution of ligands by exponential enrichment with degeneracy) procedure was utilized to isolate AlgZ targets from a random pool of DNA fragments. dSELEX is a process that allows pieces of DNA that bind with high-affinity to a protein to dominate a random DNA pool. The isolated DNA targets were cloned into a plasmid, and transformed into *E. coli* cells. Plasmids containing an AlgZ binding insert were isolated from successful clones. These clones will be examined via sequence analysis and subsequent sequence alignment should reveal any AlgZ-binding motif(s).